OIPE GRATA

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SEQUENCE LISTING

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<110>
          Xiao, Yingxao
          Feng, Xin-Hua
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          Downstream reverse primer used to amplify the upstream
          promoter containing Box D in the Human 5S RNA gene.
          sequence contains a Pst1 site at 7 bp upstream of the
          transcription site.
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25

<210> 3

<211> 269

<212> DNA

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<221> promoter

<223> Calculated BamHI-PstI fragment of the upstream promoter containing Box D in the Human 5S gene. Cloned into pBluescript-KS to give plasmid pPPVI.

<400> 3
ggatccaaaa cgctgcctcc gcgacagggc ggaggacgga gggcgtccca ggatcgtggg 60
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ctcttggggc agccaggcgc ctccttcag 269

<210> 4

<211> 84

<212> DNA

<213> Human

<220>

<221> terminator

<223> Comprises Box A, C and terminator of the human 5S RNA gene. Serves as a top strand to anneal with SEQ ID NO: 5 to create a double-stranded DNA molecule.

<400> 4

agaagacgaa gctaagcagg gtcgggcctg gttagtactt ggatgggaga ccgcctggga 60 ataccgggtg ctgtaggctt tttg 84

<210> 5

<211> 88

<212> DNA

<213> Human

<220>

<221> terminator

<223> Comprises Box A, C and terminator of the human 5S RNA gene. Serves as a top strand to anneal with SEQ ID NO: 4 to create a double-stranded DNA molecule.

<400> 5
tcgacaaaaa gcctacagca cccggtattc ccaggcggtc tcccatccaa gtactaacca 60
ggcccgaccc tgcttagctt cgtcttct 88

<210> 6

<211> 367

<212> DNA

<213> Human

<220>

<221> promoter

<223> A BamHI-SalI fragment of plasmid pPPV2 containing the upstream promoter containing Box D, A, C and the terminator of the Human 5S gene.

<400> 6
ggatccaaaa cgctgcctcc gcgacagggc ggaggacgga gggcgtccca ggatcgtggg 60
ccctgggcct gacgcctcgg agcactccct gctccgagcg ggcccgatgt ggtggaagct 120
cgggagcgcg ggagccgggg gaaggccgcg ggcagccgtc gggggtcccc gatccgagcc 180
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ctcttggggc agccaggcgc ctccttcagg aattcgatag aagacgaagc taagcagggt 300
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tgtcgac 367

<210> 7

<211> 51

<212> DNA

<213> Human

<220>

<221> misc_RNA

- <222> Positioned with PstI at the 5' end and BbsI at the 3' end.
- Contains designed siRNA sequence. Serves as a top strand to anneal with SEQ ID NO: 8 to create a double-stranded DNA molecule. The "n" bases represent any of the a, g, c, or t bases.

<400> 7

gennnnnnn nnnnnnnnn nttteggnnn nnnnnnnnn nnnnnntttt t 51

- <210> 8
- <211> 59
- <212> DNA
- <213> Human
- <220>
- <221> misc RNA
- <222> Positioned with PstI at the 5' end and BbsI at the 3' end.
- Contains designed siRNA sequence. Serves as a top strand to anneal with SEQ ID NO: 7 to create a double-stranded DNA molecule. The "n" bases represent any of the a, g, c, or t bases.

<400> 8

agctaaaaan nnnnnnnnn nnnnnnnncc gaaannnnnn nnnnnnnnn nnngctgca 59

- <210> 9
- <211> 399
- <212> DNA
- <213> Human
- <220>
- <221> misc structure
- <222> A BamHI-SalI fragment of plasmid pPPV2 containing the siRNA
 design.
- The second stretch of the 19 "n" bases are complementary and reverse to the first stretch. The "n" bases represent any of the a, g, c, or t bases.

<400> 9

ggatccaaaa cgctgcctcc gcgacagggc ggaggacgga gggcgtccca ggatcgtggg 60 ccctgggcct gacgcctcgg agcactccct gctccgagcg ggcccgatgt ggtggaagct 120 cgggagcgcg ggagccggg gaaggccgcg ggcagccgtc gggggtcccc gatccgagcc 180 ccgcggcccc gggctggcgg tgtcggctgc aatccggcgg gcacggccgg ccgggctggg 240

ctcttggggc agccaggcgc ctccttcagc nnnnnnnnn nnnnnnnnn ttcggnnnnn 300 nnnnnnnnnn nnnnttttta gctaagcagg gtcgggcctg gttagtactt ggatgggaga 360 ccgcctggga ataccgggtg ctgtaggctt tttgtcgac 399